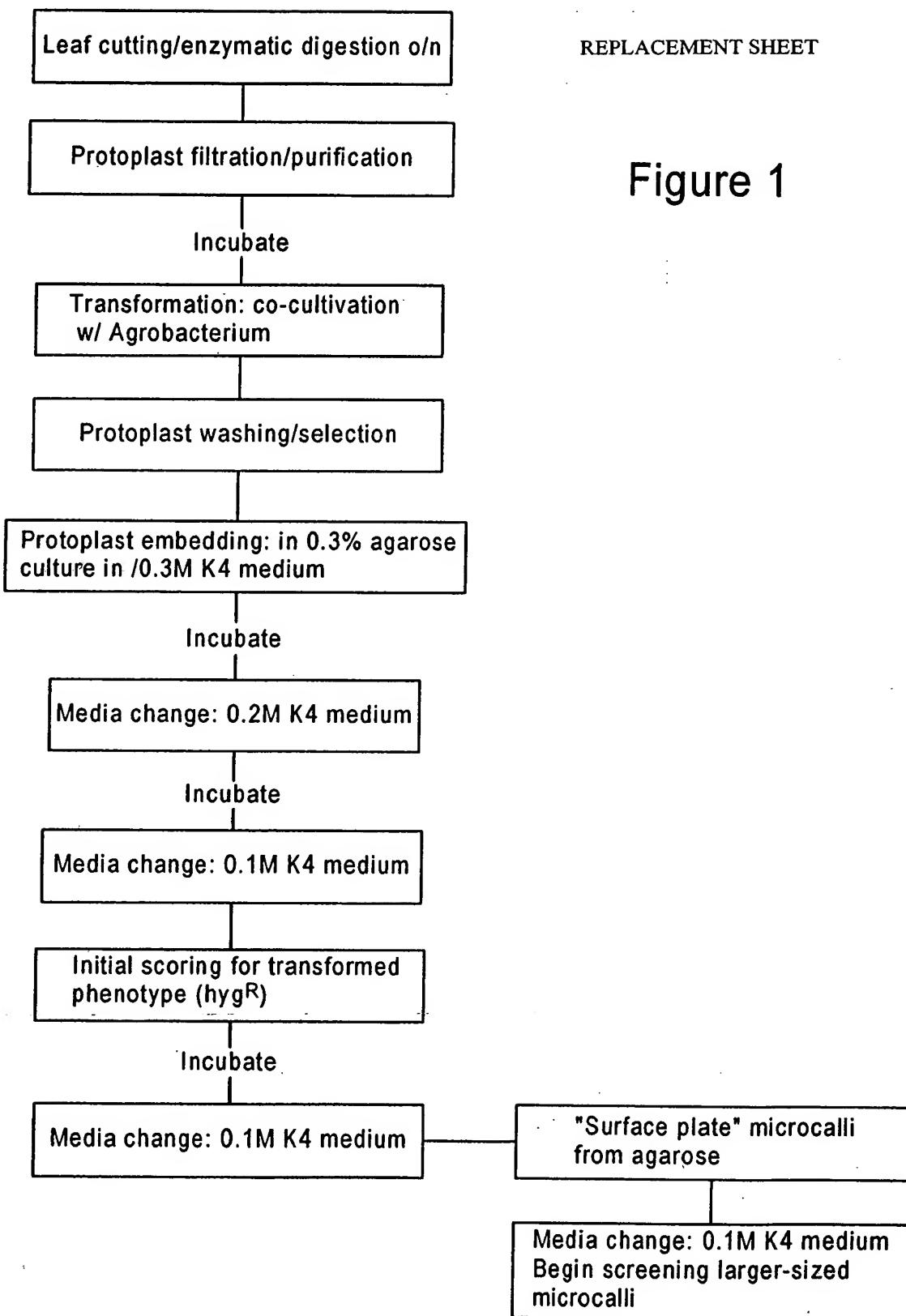
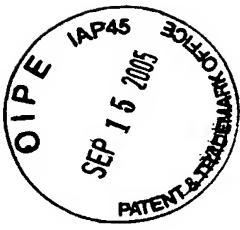




SN 10/045,677
filed 1/15/02





REPLACEMENT SHEET

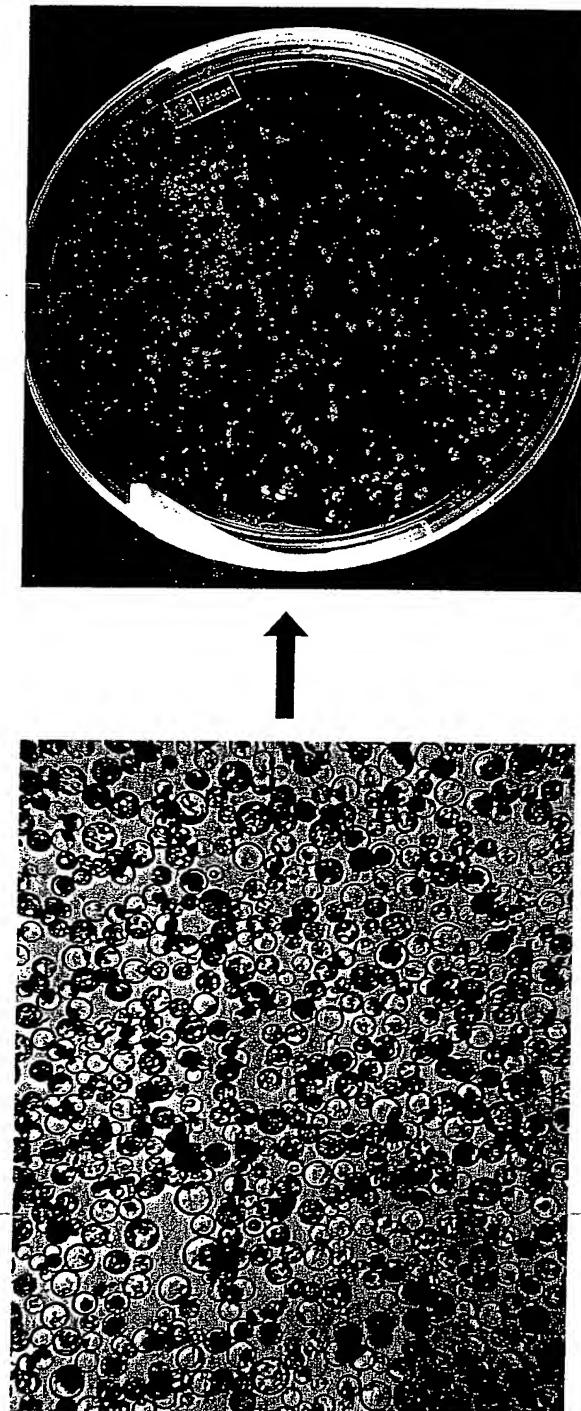
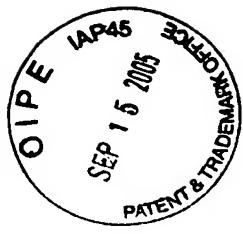


Figure 2A

Figure 2B

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REPLACEMENT SHEET

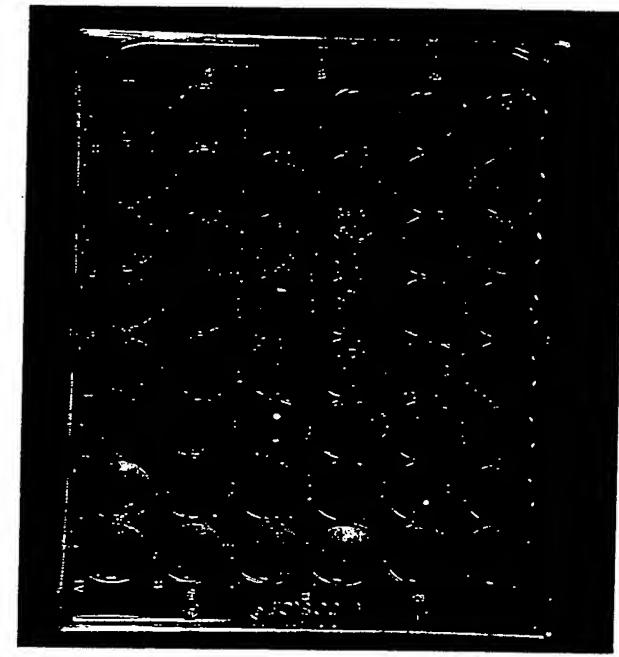


Figure 3B

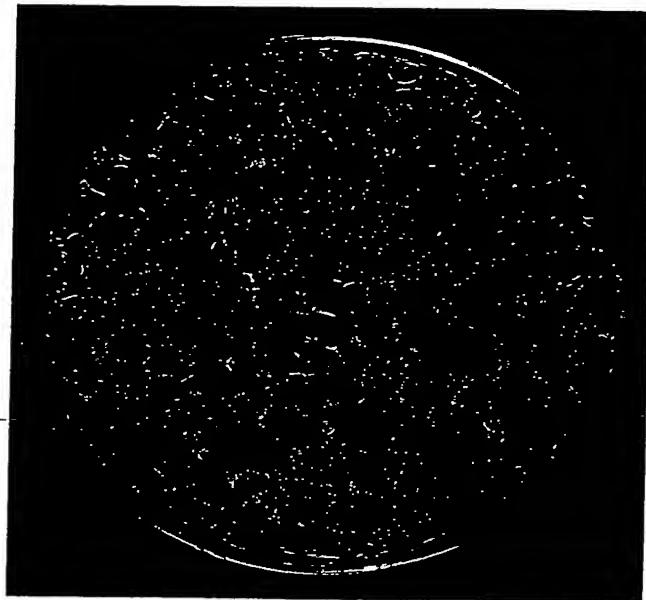
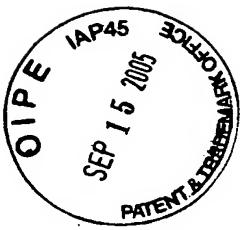


Figure 3A

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REPLACEMENT SHEET

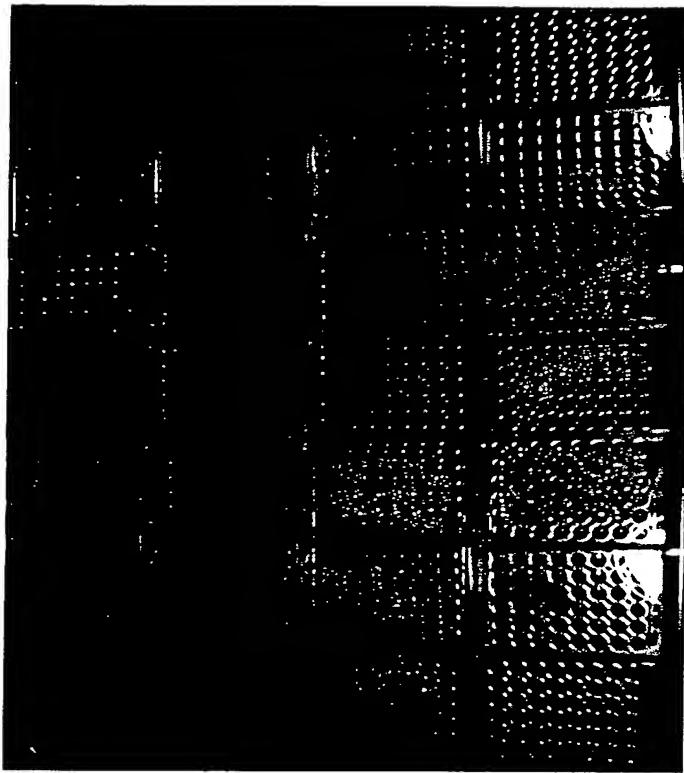
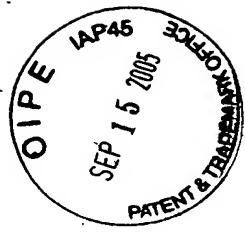


Figure 4

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REPLACEMENT SHEET

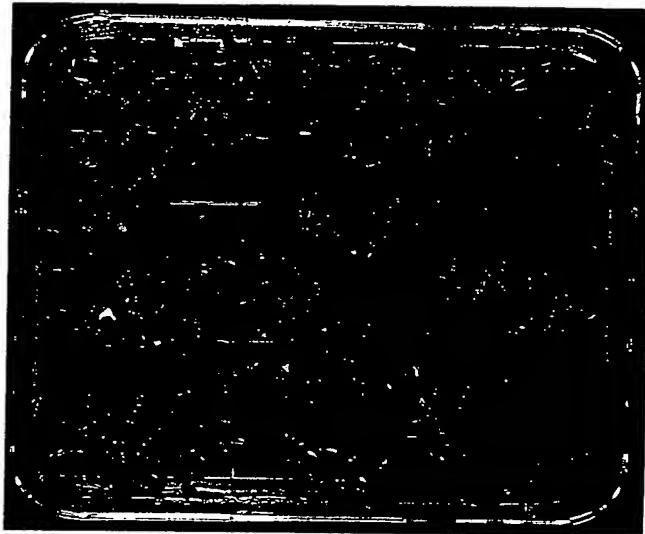


Figure 5B

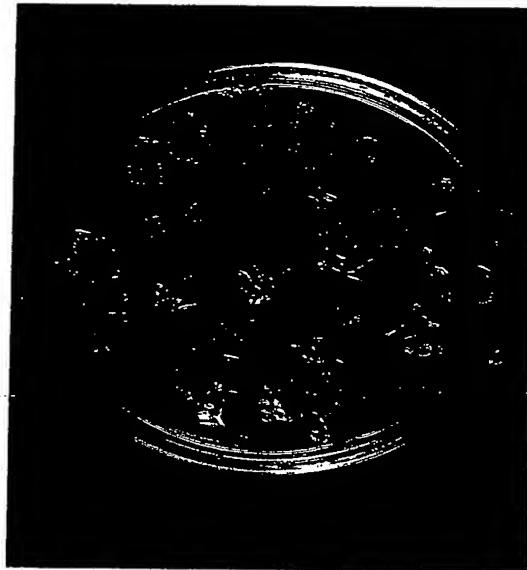


Figure 5A

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REPLACEMENT SHEET

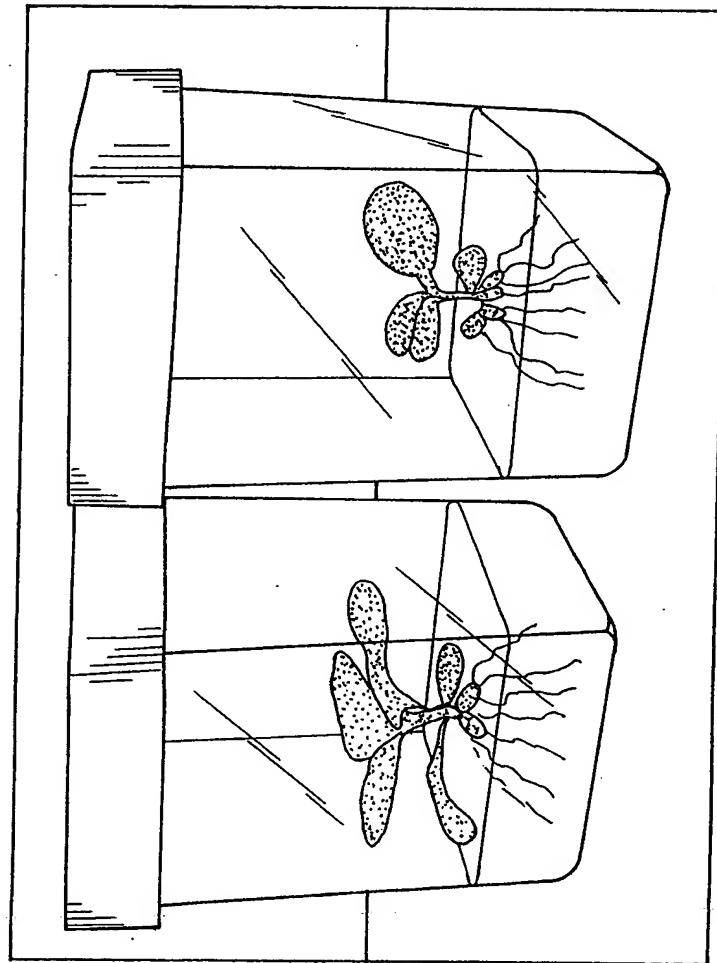
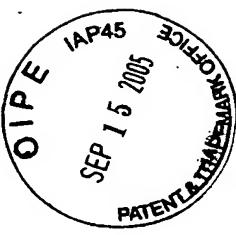


Figure 6



REPLACEMENT SHEET

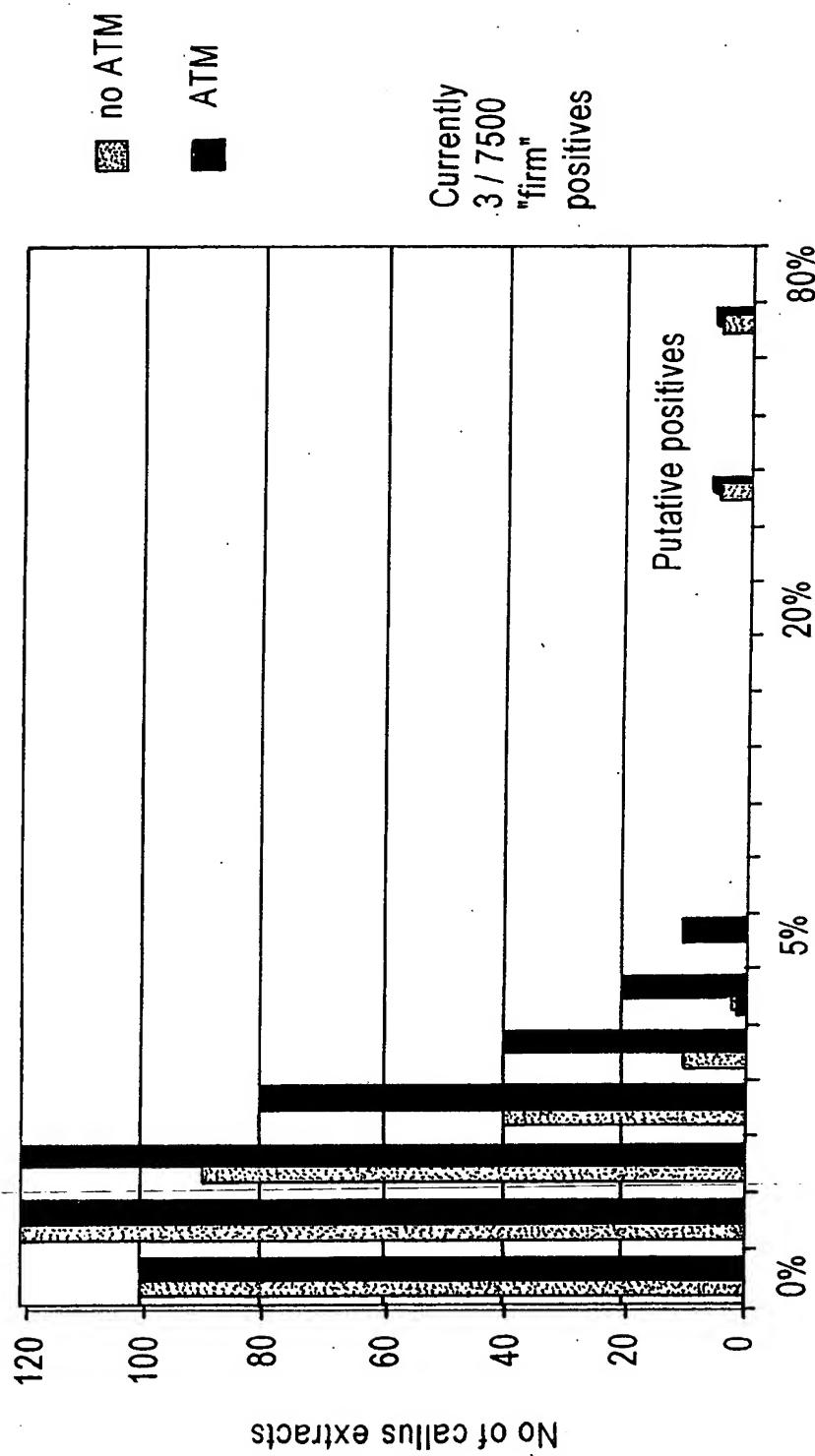


Figure 7

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REPLACEMENT SHEET

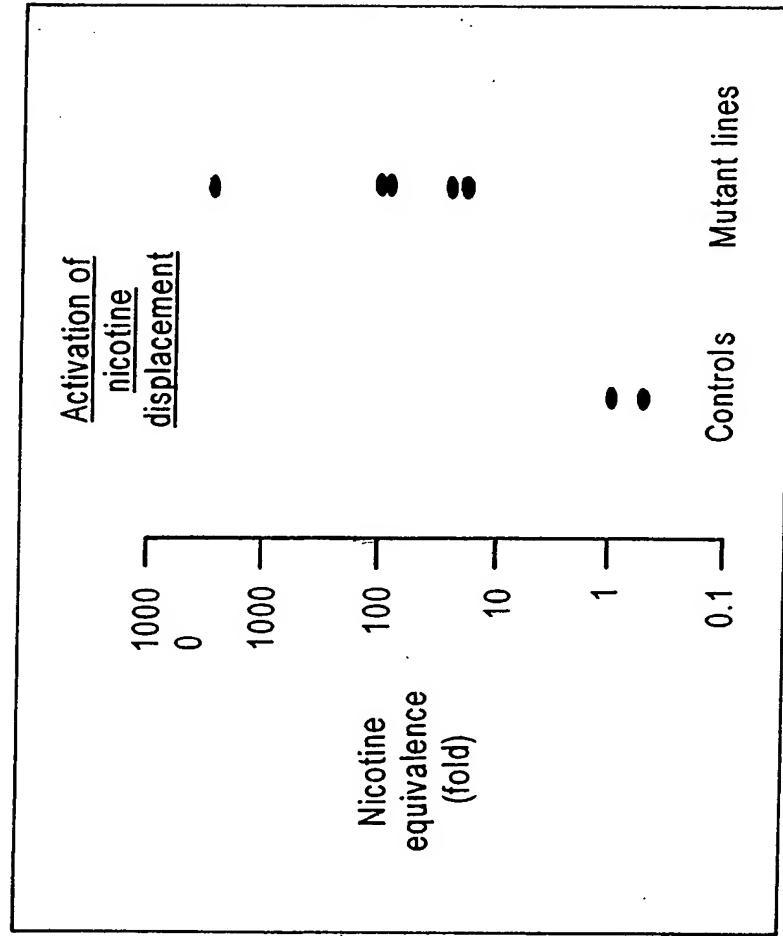
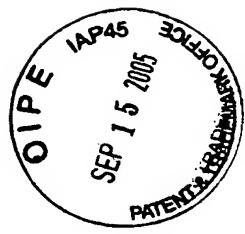


Figure 8



REPLACEMENT SHEET

[3 H]-nicotine binding displacement

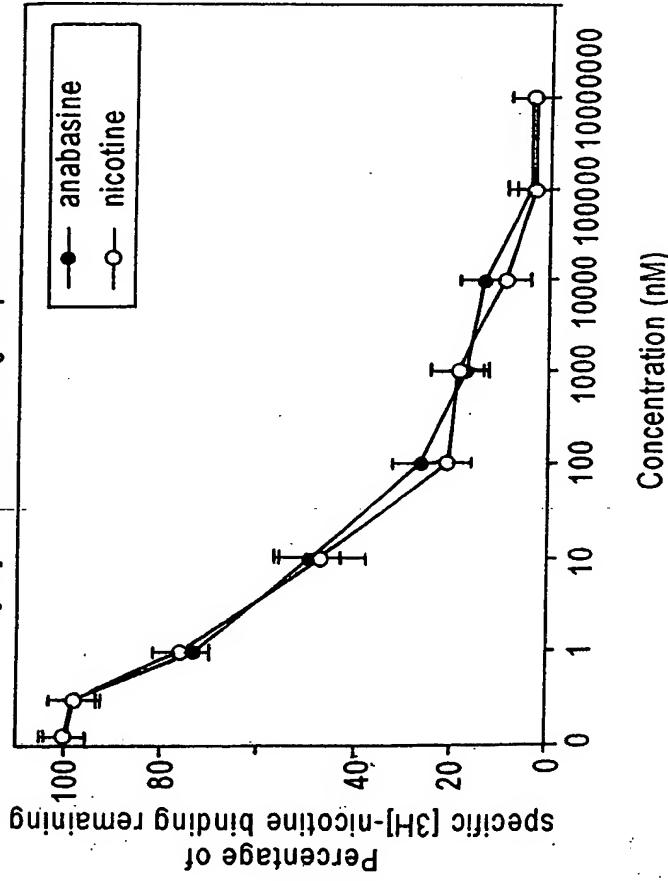


Figure 9A

[3 H]-epibatidine binding displacement

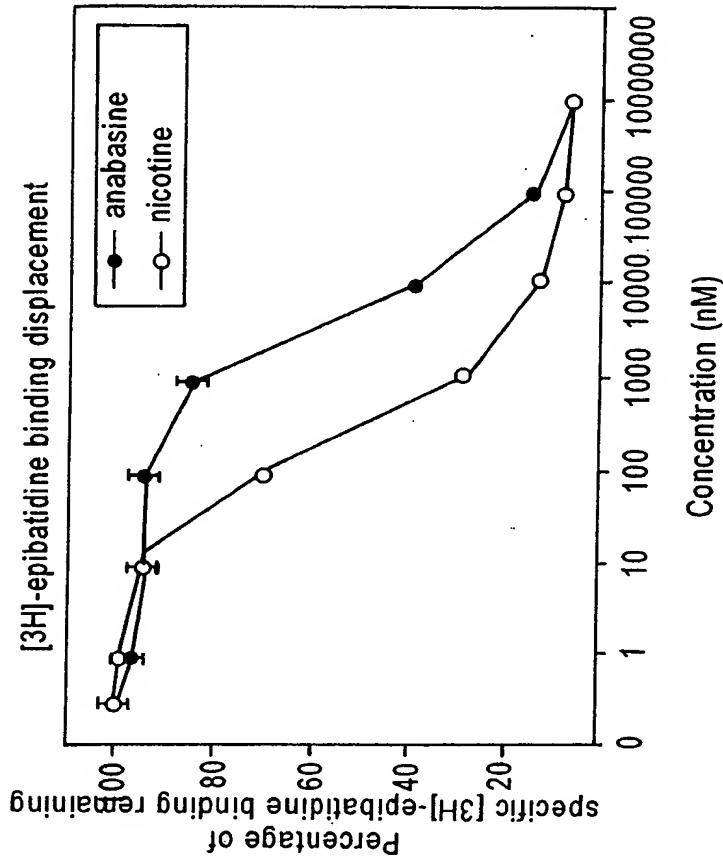
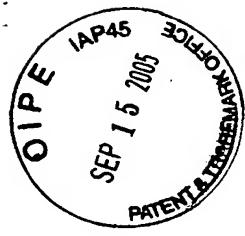


Figure 9B



REPLACEMENT SHEET

[3 H]-nicotine binding displacement

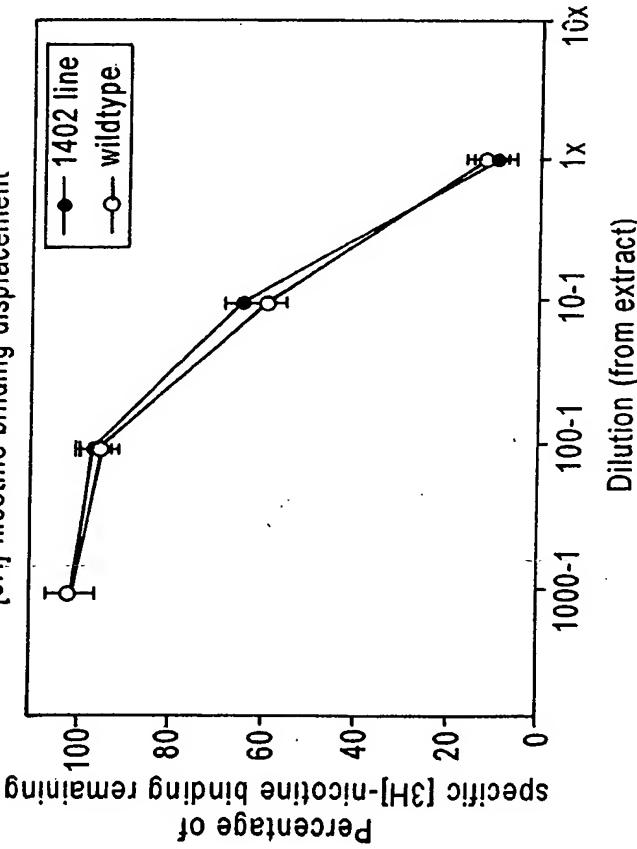


Figure 9C

[3 H]-epibatidine binding displacement

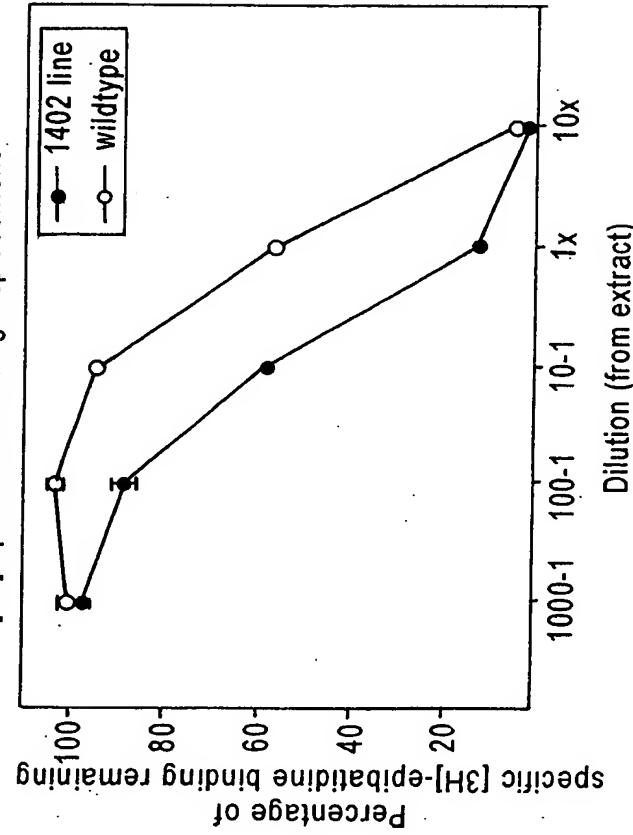
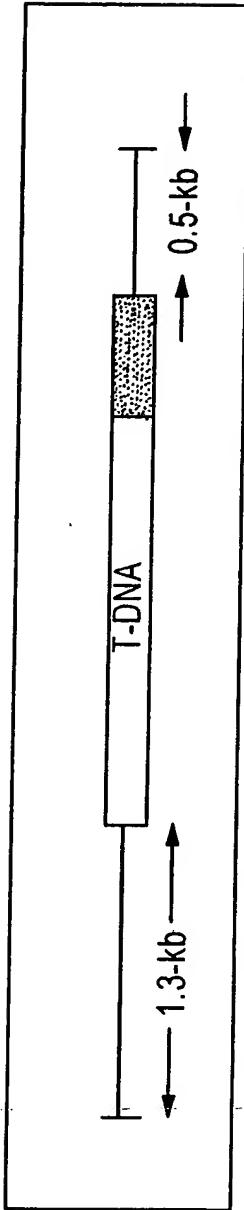


Figure 9D



REPLACEMENT SHEET

- * Genomic DNA adjacent to the activation-tagging element has been cloned from using plasmid rescue.



- * 1.8-kb of plant genomic DNA has been recovered. Initial sequence analysis indicates homology with unknown expressed sequenced tags in soybean and barley roots.
- * More extensive analysis is now underway using the cloned fragments as starting material (e.g., library probing, Northern analysis, etc)

Figure 10